

ACCESSION NUMBER: 2006:710890 CAPLUS  
 DOCUMENT NUMBER: 145:161779  
 TITLE: Fibrous protein fusions with mineralization domains and use in the formation of advanced organic/inorganic composite materials  
 INVENTOR(S): Kaplan, David L.; Huang, Jia; Wong Po Foo, Cheryl; Naik, Rajesh; George, Anne  
 PATENT ASSIGNEE(S): Trustees of Tufts College, USA; United States of America as Represented by the Secretary of the Air Force; The Board of Trustees of the University of Illinois  
 SOURCE: PCT Int. Appl., 55 pp.  
 CODEN: PIXXD2  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 1  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2006076711	A2	20060720	WO 2006-US1536	20060117
W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KM, KN, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, LY, MA, MD, MG, MK, MN, MW, MX, MZ, NA, NG, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SM, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW RW: AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IS, IT, LT, LU, LV, MC, NL, PL, PT, RO, SE, SI, SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG, BW, GH, GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				

PRIORITY APPLN. INFO.: US 2005-644264P P 20050114

AB The claimed invention provides a fusion polypeptide comprising a fibrous protein domain and a mineralization domain, and the resulting fusion protein used to form an organic-inorg. composite. Thus, the R5 peptide (SSKKS GSYSGSKGSKRRIL) of silaffin-1 from *Cylindrotheca fusiformis* is genetically fused to a 15-mer of the consensus repeat unit (SGRGGLGGQGAGAAAAAGGAGQGGYGGGLGSQGT with CRGD linker) of spidroin 1 from the golden orb spider *Nephila clavipes*. The purpose of fusing this silicification-inducing peptide unit to genetically engineered silk is to combine the properties of the silk whether in the form of films or other such spun fibers to the silica-precipitating properties of R5 under ambient conditions to produce biomaterials with controlled silica morphologies on the surface. Silicification reactions using tetraethoxysilane on synthetic spider silk protein films yielded spherical silica structures with diams. ranging from .apprx.0.5-2.0  $\mu$ m only when the silica precipitating domain, R5 peptide, was fused to the C-terminus of the silk proteins. These organic-inorg. composites can be constructed from the nano- to the macro-scale depending on the size of the fibrous protein fusion domain used. The composites can also be loaded with other compds. (e.g., dyes, drugs, enzymes) depending on the goal for the materials, to further enhance function. This can be achieved during assembly of the material or during the mineralization step in materials formation.

L6 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2005:394700 CAPLUS  
 DOCUMENT NUMBER: 142:428879  
 TITLE: Entrapment of biomolecules and inorganic nanoparticles by biosilicification  
 INVENTOR(S): Naik, Rajesh R.; Stone, Morley O.; Spain, Jim C.; Luckarift, Heather R.

PATENT ASSIGNEE(S): USA  
 SOURCE: U.S. Pat. Appl. Publ., 16 pp.  
 CODEN: USXXCO  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 1  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 2005095690	A1	20050505	US 2004-803633	20040318
PRIORITY APPLN. INFO.:			US 2003-517227P	P 20031031

AB A method of immobilizing  $\geq 1$  mol. in a silica matrix to form a biosilicification product. The mol. may be immobilized in the silica matrix at substantially the same time as the silica matrix is formed. The method comprises combining  $\geq 1$  silaffin polypeptide,  $\geq 1$  mol., and  $\geq 1$  hydroxylated water-soluble derivative to form the biosilicification product.

The silaffin polypeptide may be Sill protein from *Cylindrotheca fusiformis*, a fragment of the Sill protein, poly-L-lysine, or a synthetic polypeptide having affinity for silica. The mol. may be an enzyme, a protein, a polypeptide, an antibody, an antigen, poly(nucleic) acids, microbial cells, plant cells, or animal cells. The hydroxylated water-soluble derivative may be silicic acid.

L6 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2007 ACS on STN  
 ACCESSION NUMBER: 2004:79316 CAPLUS  
 DOCUMENT NUMBER: 140:402326  
 TITLE: Enzyme immobilization in a biomimetic silica support  
 AUTHOR(S): Luckarift, Heather R.; Spain, Jim C.; Naik, Rajesh R.; Stone, Morley O.  
 CORPORATE SOURCE: Air Force Research Laboratory, Airbase Technologies Division, Suite #2, Tyndall Air Force Base, FL, 32403-5323, USA  
 SOURCE: Nature Biotechnology (2004), 22(2), 211-213  
 CODEN: NABIF9; ISSN: 1087-0156  
 PUBLISHER: Nature Publishing Group  
 DOCUMENT TYPE: Journal  
 LANGUAGE: English

AB Robust immobilization techniques that preserve the activity of biomols. have many potential applications. Silicates, primarily in the form of sol-gel composites or functionalized mesoporous silica, have been used to encapsulate a wide variety of biomols. but the harsh conditions required for chemical synthesis limit their applicability. Silaffin polypeptides from diatoms catalyze the formation of silica in vitro at neutral pH and ambient temperature and pressure. Here we show that butyrylcholinesterase entrapped during the precipitation of silica nanospheres retained all of its activity. Ninety percent of the soluble enzyme was immobilized, and the immobilized enzyme was substantially more stable than the free enzyme. The mech. properties of silica nanospheres facilitated application in a flow-through reactor. The use of biosilica for enzyme immobilization combines the excellent support properties of a silica matrix with a benign immobilization method that retains enzyme activity.

REFERENCE COUNT: 18 THERE ARE 18 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> d ibib abs 17 1-14

L7 ANSWER 1 OF 14 MEDLINE on STN  
 ACCESSION NUMBER: 2004496677 MEDLINE  
 DOCUMENT NUMBER: PubMed ID: 15304518  
 TITLE: Silica morphogenesis by alternative processing of silaffins in the diatom *Thalassiosira pseudonana*.  
 AUTHOR: Poulsen Nicole; Kroger Nils  
 CORPORATE SOURCE: Lehrstuhl Biochemie I, Universitätsstr. 31, Universität Regensburg, 93053 Regensburg, Germany.  
 SOURCE: The Journal of biological chemistry, (2004 Oct 8) Vol. 279, No. 41, pp. 42993-9. Electronic Publication: 2004-08-10. Journal code: 2985121R. ISSN: 0021-9258.  
 PUB. COUNTRY: United States  
 DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE) (RESEARCH SUPPORT, NON-U.S. GOV'T)  
 LANGUAGE: English  
 FILE SEGMENT: Priority Journals  
 OTHER SOURCE: GENBANK-AY706749; GENBANK-AY706750; GENBANK-AY706751  
 ENTRY MONTH: 200411  
 ENTRY DATE: Entered STN: 7 Oct 2004  
 Last Updated on STN: 19 Dec 2004  
 Entered Medline: 24 Nov 2004

AB For almost 200 years scientists have been fascinated by the ornate cell walls of the diatoms. These structures are made of amorphous silica, exhibiting species-specific, mostly porous patterns in the nano- to micrometer range. Recently, from the diatom *Cylindrotheca fusiformis* unusual phosphoproteins (termed silaffins) and long chain polyamines have been identified and implicated in biosilica formation. However, analysis of the role of silaffins in morphogenesis of species-specific silica structures has so far been hampered by the difficulty of obtaining structural data from these extremely complex proteins. In the present study, the five major silaffins from the diatom *Thalassiosira pseudonana* (tpSil1H, -1L, -2H, -2L, and -3) have been isolated, functionally analyzed, and structurally characterized, making use of the recently available genome data from this organism. Surprisingly, the silaffins of *T. pseudonana* and *C. fusiformis* share no sequence homology but are similar regarding amino acid composition and post-translational modifications. Silaffins tpSil1H and -2H are higher molecular mass isoforms of tpSil1L and -2L, respectively, generated in vivo by alternative processing of the same precursor polypeptides. Interestingly, only tpSil1H and -2H but not tpSil1L and -2L induce the formation of porous silica patterns in vitro, suggesting that the alternative processing event is an important step in morphogenesis of *T. pseudonana* biosilica.

L7 ANSWER 2 OF 14 BIOSIS COPYRIGHT (c) 2007 The Thomson Corporation on STN  
 ACCESSION NUMBER: 2005:14060 BIOSIS  
 DOCUMENT NUMBER: PREV200500015265  
 TITLE: Silica morphogenesis by alternative processing of silaffins in the diatom *Thalassiosira pseudonana*.  
 AUTHOR(S): Poulsen, Nicole; Kroeger, Nils [Reprint Author]  
 CORPORATE SOURCE: Lehrstuhl Biochem 1, Univ Regensburg, Univ Str 31, D-93053, Regensburg, Germany  
 nils.kroeger@vkl.uni-regensburg.de  
 SOURCE: Journal of Biological Chemistry, (October 8 2004) Vol. 279, No. 41, pp. 42993-42999, 42984. print. CODEN: JBCHA3. ISSN: 0021-9258.  
 DOCUMENT TYPE: Article  
 LANGUAGE: English  
 ENTRY DATE: Entered STN: 22 Dec 2004  
 Last Updated on STN: 22 Dec 2004

AB For almost 200 years scientists have been fascinated by the ornate cell walls of the diatoms. These structures are made of amorphous silica, exhibiting species-specific, mostly porous patterns in the

nano- to micrometer range. Recently, from the diatom *Cylindrotheca fusiformis* unusual phosphoproteins ( termed silaffins) and long chain polyamines have been identified and implicated in biosilica formation. However, analysis of the role of silaffins in morphogenesis of species-specific silica structures has so far been hampered by the difficulty of obtaining structural data from these extremely complex proteins. In the present study, the five major silaffins from the diatom *Thalassiosira pseudonana* (tpSillH, - 1L, - 2H, - 2L, and - 3) have been isolated, functionally analyzed, and structurally characterized, making use of the recently available genome data from this organism. Surprisingly, the silaffins of *T. pseudonana* and *C. fusiformis* share no sequence homology but are similar regarding amino acid composition and posttranslational modifications. Silaffins tpSillH and - 2H are higher molecular mass isoforms of tpSillL and - 2L, respectively, generated in vivo by alternative processing of the same precursor polypeptides. Interestingly, only tpSillH and - 2H but not tpSillL and - 2L induce the formation of porous silica patterns in vitro, suggesting that the alternative processing event is an important step in morphogenesis of *T. pseudonana* biosilica.

L7 ANSWER 3 OF 14 BIOSIS COPYRIGHT (c) 2007 The Thomson Corporation on STN  
 ACCESSION NUMBER: 2002:508051 BIOSIS  
 DOCUMENT NUMBER: PREV200200508051  
 TITLE: Synthesis of silica nanostructures at neutral pH using catalytic polypeptides.  
 AUTHOR(S): Clarson, Stephen J. [Reprint author]; Whitlock, Patrick William; Patwardhan, Siddharth V. [Reprint author]; Brott, Lawrence L.; Naik, Rajesh R.; Stone, Morley O.  
 CORPORATE SOURCE: Department of Materials Science and Engineering, University of Cincinnati, 492 Rhodes Hall, Cincinnati, OH, 45221-0012, USA  
 sclarson@uceng.uc.edu  
 SOURCE: Abstracts of Papers American Chemical Society, (2002) Vol. 223, No. 1-2, pp. PMSE 231. print.  
 Meeting Info.: 223rd National Meeting of the American Chemical Society. Orlando, FL, USA. April 07-11, 2002.  
 CODEN: ACSRAL. ISSN: 0065-7727.  
 DOCUMENT TYPE: Conference; (Meeting)  
 Conference; Abstract; (Meeting Abstract)  
 LANGUAGE: English  
 ENTRY DATE: Entered STN: 2 Oct 2002  
 Last Updated on STN: 2 Oct 2002

L7 ANSWER 4 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN  
 ACCESSION NUMBER: 2006:710890 CAPLUS  
 DOCUMENT NUMBER: 145:161779  
 TITLE: Fibrous protein fusions with mineralization domains and use in the formation of advanced organic/inorganic composite materials  
 INVENTOR(S): Kaplan, David L.; Huang, Jia; Wong Po Foo, Cheryl; Naik, Rajesh; George, Anne  
 PATENT ASSIGNEE(S): Trustees of Tufts College, USA; United States of America as Represented by the Secretary of the Air Force; The Board of Trustees of the University of Illinois  
 SOURCE: PCT Int. Appl., 55 pp.  
 CODEN: PIXXD2  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 1  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
------------	------	------	-----------------	------

-----  
 WO 2006076711                      A2                      20060720                      WO 2006-US1536                      20060117  
 W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH,  
 CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD,  
 GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KM, KN, KP, KR,  
 KZ, LC, LK, LR, LS, LT, LU, LV, LY, MA, MD, MG, MK, MN, MW, MX,  
 MZ, NA, NG, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE,  
 SG, SK, SL, SM, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC,  
 VN, YU, ZA, ZM, ZW  
 RW: AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE,  
 IS, IT, LT, LU, LV, MC, NL, PL, PT, RO, SE, SI, SK, TR, BF, BJ,  
 CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG, BW, GH,  
 GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY,  
 KG, KZ, MD, RU, TJ, TM

PRIORITY APPLN. INFO.:                      US 2005-644264P                      P 20050114

AB The claimed invention provides a fusion polypeptide comprising a fibrous protein domain and a mineralization domain, and the resulting fusion protein used to form an organic-inorg. composite. Thus, the R5 peptide (SSKKSGSYSGSKGSKRRIL) of silaffin-1 from *Cylindrotheca fusiformis* is genetically fused to a 15-mer of the consensus repeat unit (SGRGGGLGGQGAGAAAAAGGAGQGGYGGGLGSQGT with CRGD linker) of spidroin 1 from the golden orb spider *Nephila clavipes*. The purpose of fusing this silicification-inducing peptide unit to genetically engineered silk is to combine the properties of the silk whether in the form of films or other such spun fibers to the silica-precipitating properties of R5 under ambient conditions to produce biomaterials with controlled silica morphologies on the surface. Silicification reactions using tetraethoxysilane on synthetic spider silk protein films yielded spherical silica structures with diams. ranging from .apprx.0.5-2.0  $\mu$ m only when the silica precipitating domain, R5 peptide, was fused to the C-terminus of the silk proteins. These organic-inorg. composites can be constructed from the nano- to the macro-scale depending on the size of the fibrous protein fusion domain used. The composites can also be loaded with other compds. (e.g., dyes, drugs, enzymes) depending on the goal for the materials, to further enhance function. This can be achieved during assembly of the material or during the mineralization step in materials formation.

L7 ANSWER 5 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN  
 ACCESSION NUMBER: 2005:394700 CAPLUS  
 DOCUMENT NUMBER: 142:428879  
 TITLE: Entrapment of biomolecules and inorganic nanoparticles by biosilicification  
 INVENTOR(S): Naik, Rajesh R.; Stone, Morley O.; Spain, Jim C.; Luckarift, Heather R.  
 PATENT ASSIGNEE(S): USA  
 SOURCE: U.S. Pat. Appl. Publ., 16 pp.  
 CODEN: USXXCO  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 1  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 2005095690	A1	20050505	US 2004-803633	20040318
PRIORITY APPLN. INFO.:			US 2003-517227P	P 20031031
AB A method of immobilizing $\geq 1$ mol. in a silica matrix to form a biosilicification product. The mol. may be immobilized in the silica matrix at substantially the same time as the silica matrix is formed. The method comprises combining $\geq 1$ silaffin polypeptide, $\geq 1$ mol., and $\geq 1$ hydroxylated water-soluble derivative to form the biosilicification product.				

The

silaffin polypeptide may be Sil1 protein from *Cylindrotheca fusiformis*, a fragment of the Sil1 protein, poly-L-lysine, or a synthetic polypeptide having affinity for silica. The mol. may be an enzyme, a protein, a polypeptide, an antibody, an antigen, poly(nucleic) acids, microbial cells, plant cells, or animal cells. The hydroxylated water-soluble derivative may be silicic acid.

L7 ANSWER 6 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2004:807360 CAPLUS  
DOCUMENT NUMBER: 142:1518  
TITLE: Silica Morphogenesis by Alternative Processing of Silaffins in the Diatom *Thalassiosira pseudonana*  
AUTHOR(S): Poulsen, Nicole; Kroeger, Nils  
CORPORATE SOURCE: Lehrstuhl Biochemie I, Universitaet Regensburg, Regensburg, 93053, Germany  
SOURCE: Journal of Biological Chemistry (2004), 279(41), 42993-42999  
CODEN: JBCHA3; ISSN: 0021-9258  
PUBLISHER: American Society for Biochemistry and Molecular Biology  
DOCUMENT TYPE: Journal  
LANGUAGE: English

AB For almost 200 yr scientists have been fascinated by the ornate cell walls of the diatoms. These structures are made of amorphous silica, exhibiting species-specific, mostly porous patterns in the nano- to micrometer range. Recently, from the diatom *Cylindrotheca fusiformis* unusual phosphoproteins (termed silaffins) and long chain polyamines have been identified and implicated in biosilica formation. However, anal. of the role of silaffins in morphogenesis of species-specific silica structures has so far been hampered by the difficulty of obtaining structural data from these extremely complex proteins. In the present study, the five major silaffins from the diatom *Thalassiosira pseudonana* (tpSil1H, -1L, -2H, -2L, and -3) have been isolated, functionally analyzed, and structurally characterized, making use of the recently available genome data from this organism. Surprisingly, the silaffins of *T. pseudonana* and *C. fusiformis* share no sequence homol. but are similar regarding amino acid composition and post-translational modifications. Silaffins tpSil1H and -2H are higher mol. mass isoforms of tpSil1L and -2L, resp., generated in vivo by alternative processing of the same precursor polypeptides. Interestingly, only tpSil1H and -2H but not tpSil1L and -2L induce the formation of porous silica patterns in vitro, suggesting that the alternative processing event is an important step in morphogenesis of *T. pseudonana* biosilica.

REFERENCE COUNT: 25 THERE ARE 25 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 7 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2004:79316 CAPLUS  
DOCUMENT NUMBER: 140:402326  
TITLE: Enzyme immobilization in a biomimetic silica support  
AUTHOR(S): Luckarift, Heather R.; Spain, Jim C.; Naik, Rajesh R.; Stone, Morley O.  
CORPORATE SOURCE: Air Force Research Laboratory, Airbase Technologies Division, Suite #2, Tyndall Air Force Base, FL, 32403-5323, USA  
SOURCE: Nature Biotechnology (2004), 22(2), 211-213  
CODEN: NABIF9; ISSN: 1087-0156  
PUBLISHER: Nature Publishing Group  
DOCUMENT TYPE: Journal  
LANGUAGE: English

AB Robust immobilization techniques that preserve the activity of biomols. have many potential applications. Silicates, primarily in the form of sol-gel composites or functionalized mesoporous silica, have been used to encapsulate a wide variety of biomols. but the harsh conditions required for chemical synthesis limit their applicability. Silaffin polypeptides from diatoms catalyze the formation of silica in vitro at neutral pH and ambient temperature and pressure. Here we show that butyrylcholinesterase entrapped during the precipitation of silica nanospheres retained all of its activity. Ninety percent of the soluble enzyme was immobilized, and the immobilized enzyme was substantially more stable than the free enzyme. The mech: properties of silica nanospheres facilitated application in a flow-through reactor. The use of biosilica for enzyme immobilization combines the excellent support properties of a silica matrix with a benign immobilization method that retains enzyme activity.

REFERENCE COUNT: 18 THERE ARE 18 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 8 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2002:896469 CAPLUS

DOCUMENT NUMBER: 138:238701

TITLE: Silicification and Biosilicification. Part 4. Effect of Template Size on the Formation of Silica

AUTHOR(S): Patwardhan, Siddharth V.; Clarson, Stephen J.

CORPORATE SOURCE: Department of Materials Science and Engineering, University of Cincinnati, Cincinnati, OH, 45221-0012, USA

SOURCE: Journal of Inorganic and Organometallic Polymers (2002), 12(3/4), 109-116

CODEN: JIOPE4; ISSN: 1053-0495

PUBLISHER: Kluwer Academic/Plenum Publishers

DOCUMENT TYPE: Journal

LANGUAGE: English

AB Silicification at neutral pH and under ambient conditions is of growing interest due to its close relationship with biosilicification. In diatoms biosilicification has been reported to occur at (or close to) neutral pH and it has been shown that protein mols. act as catalysts/templates/scaffolds for this elegant materials chemical. In this investigation various catalysts/templates have been studied for their role in silicification in vitro. We have used functionalized C60 fullerene, R5 (an important polypeptide from the amino acid sequence of a silaffin protein), poly-L-lysine (PLL) and two poly(allylamine hydrochloride) (PAH) samples having different mol. wts. An aqueous silica precursor was used and ordered silica structures were produced in each of the systems studied. The sizes of the silica structures appear to correlate with the size, in solution, of the templating/scaffolding agents. Biol. systems exhibit hierarchical structures with remarkable control of morphologies over different length scales. The use of templating/scaffolding agents having different sizes and shapes is one possible paradigm for the production of such structures in vivo.

REFERENCE COUNT: 18 THERE ARE 18 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 9 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2002:232433 CAPLUS

TITLE: Synthesis of silica nanostructures at neutral Ph using catalytic polypeptides

AUTHOR(S): Clarson, Stephen J.; Whitlock, Patrick W.; Patwardhan, Siddharth V.; Brot, Lawrence L.; Naik, Rajesh R.; Stone, Morley O.

CORPORATE SOURCE: Department of Materials Science and Engineering, University of Cincinnati, Cincinnati, OH, 45221-0012, USA

SOURCE: PMSE Preprints (2002), 86, 81  
CODEN: PPMRA9; ISSN: 1550-6703  
PUBLISHER: American Chemical Society  
DOCUMENT TYPE: Journal; (computer optical disk)  
LANGUAGE: English

AB Marine diatoms and sponges are capable of forming beautiful silica nanostructures in vivo. Silaffin proteins have recently been isolated from the marine diatom *Cylindrotheca fusiformis* and have been shown to generate silica spheres when added to solns. of silicic acid in vitro. We report here a variety of silica structures and inorg.-organic hybrid materials that have been prepared using synthetic peptide sequences derived from the Sill gene of *Cylindrotheca fusiformis*. We are also exploring synthetic polymers that can mimic the catalytic/templating function of these biol. derived systems.

REFERENCE COUNT: 4 THERE ARE 4 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 10 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2002:191815 CAPLUS

TITLE: Synthesis of silica nanostructures at neutral pH using catalytic polypeptides

AUTHOR(S): Clarson, Stephen J.; Whitlock, Patrick William; Patwardhan, Siddharth V.; Brott, Lawrence L.; Naik, Rajesh R.; Stone, Morley O.

CORPORATE SOURCE: Department of Materials Science and Engineering, University of Cincinnati, Cincinnati, OH, 45221-0012, USA

SOURCE: Abstracts of Papers, 223rd ACS National Meeting, Orlando, FL, United States, April 7-11, 2002 (2002), PMSE-231. American Chemical Society: Washington, D. C.

CODEN: 69CKQP

DOCUMENT TYPE: Conference; Meeting Abstract

LANGUAGE: English

AB Marine diatoms and sponges are capable of forming beautiful silica nanostructures in vivo. Silaffin proteins have recently been isolated from the marine diatom *Cylindrotheca fusiformis* and have been shown to generate silica spheres when added to solns. of silicic acid in vitro. We report here a variety of silica structures and inorg.-organic hybrid materials that have been prepared using synthetic peptide sequences derived from the Sill gene of *Cylindrotheca fusiformis*. We are also exploring synthetic polymers that can mimic the catalytic / templating function of these biol. derived systems.

L7 ANSWER 11 OF 14 CAPLUS COPYRIGHT 2007 ACS on STN

ACCESSION NUMBER: 2001:200992 CAPLUS

TITLE: Biocatalysis of silica nanostructures

AUTHOR(S): Whitlock, Patrick W.; Brott, Lawrence L.; Clarson, Stephen J.; Naik, Rajesh R.; Stone, Morley O.

CORPORATE SOURCE: Materials Science and Engineering, The Polymer Research Group / Materials and Manufacturing Directorate, University of Cincinnati / Air Force Research Laboratory, Cincinnati, OH, 45221-0018, USA

SOURCE: Abstracts of Papers, 221st ACS National Meeting, San Diego, CA, United States, April 1-5, 2001 (2001) IEC-321

CODEN: 69FZD4

PUBLISHER: American Chemical Society

DOCUMENT TYPE: Journal; Meeting Abstract

LANGUAGE: English

AB Numerous examples of nanopatterning and nanostructure are commonly found in nature, most apparent in the marine diatoms and sponges. In vivo biosilification allows these organisms to control structural morphol. at the nanometer level. Understanding how nature performs this exquisite



control and duplicating this process has numerous applications in materials science. Silaffins, a set of cationic polypeptides isolated from the diatom *Cylindrotheca fusiformis*, can generate a network of silica nanospheres when added to a solution of silicic acid in vitro. Using a short synthetic peptide derived from the Silaffin 1 (Sill) protein of *C. fusiformis*, we produced a variety of silica nanostructures. The produced structures range in morphol. from common spheres to highly organized and complex fibrillar geometries that display remarkable organization at the nanometer size-scale. We are currently investigating the mol. orientation present in these morphologies and developing new methods to control the deposition of silica for nanoapplications.

L7 ANSWER 12 OF 14 LIFESCI COPYRIGHT 2007 CSA on STN  
ACCESSION NUMBER: 2005:92539 LIFESCI  
TITLE: Silica Morphogenesis by Alternative Processing of Silaffins in the Diatom *Thalassiosira pseudonana*  
AUTHOR: Poulsen, Nicole; Kroeger, Nils  
CORPORATE SOURCE: Lehrstuhl Biochemie I, Universitaetsstr. 31, Universitaet Regensburg, 93053 Regensburg, Germany  
SOURCE: Journal of Biological Chemistry [J. Biol. Chem.], (2004)1008  
) vol. 279, no. 41, pp. 42993-42999.  
ISSN: 0021-9258.  
DOCUMENT TYPE: Journal  
FILE SEGMENT: K  
LANGUAGE: English  
SUMMARY LANGUAGE: English

AB For almost 200 years scientists have been fascinated by the ornate cell walls of the diatoms. These structures are made of amorphous silica, exhibiting species-specific, mostly porous patterns in the nano- to micrometer range. Recently, from the diatom *Cylindrotheca fusiformis* unusual phosphoproteins (termed silaffins) and long chain polyamines have been identified and implicated in biosilica formation. However, analysis of the role of silaffins in morphogenesis of species-specific silica structures has so far been hampered by the difficulty of obtaining structural data from these extremely complex proteins. In the present study, the five major silaffins from the diatom *Thalassiosira pseudonana* (tpSillH, -1L, -2H, -2L, and -3) have been isolated, functionally analyzed, and structurally characterized, making use of the recently available genome data from this organism. Surprisingly, the silaffins of *T. pseudonana* and *C. fusiformis* share no sequence homology but are similar regarding amino acid composition and post-translational modifications. Silaffins tpSillH and -2H are higher molecular mass isoforms of tpSillL and -2L, respectively, generated in vivo by alternative processing of the same precursor polypeptides. Interestingly, only tpSillH and -2H but not tpSillL and -2L induce the formation of porous silica patterns in vitro, suggesting that the alternative processing event is an important step in morphogenesis of *T. pseudonana* biosilica.

L7 ANSWER 13 OF 14 EMBASE COPYRIGHT (c) 2007 Elsevier B.V. All rights reserved on STN  
ACCESSION NUMBER: 2005168410 EMBASE  
TITLE: Prospects in diatom research.  
AUTHOR: Lopez P.J.; Descles J.; Allen A.E.; Bowler C.  
CORPORATE SOURCE: P.J. Lopez, CNRS FRE-2910 Signalisation M., Ecole Normale Supérieure, 46 rue d'Ulm, 75005 Paris, France.  
pjlopez@biologie.ens.fr  
SOURCE: Current Opinion in Biotechnology, (2005) Vol. 16, No. 2, pp. 180-186. .  
Refs: 53  
ISSN: 0958-1669 CODEN: CUOBE3  
COUNTRY: United Kingdom

DOCUMENT TYPE: Journal; General Review  
FILE SEGMENT: 004 Microbiology  
LANGUAGE: English  
SUMMARY LANGUAGE: English  
ENTRY DATE: Entered STN: 12 May 2005  
Last Updated on STN: 12 May 2005

AB Diatoms are unicellular photosynthetic eukaryotes that play a major role in the global cycling of carbon and silicon. They are believed to have arisen from a secondary endosymbiotic event between two eukaryotes, a red alga and a flagellated heterotroph. Recent analysis of a diatom genome indeed reveals a 'mosaic' nature, with genes derived from plant, animal and bacterial lineages. Advances in molecular genomics are facilitating the use of diatom-specific genes or pathways for biotechnology. Another interest is in understanding the artistry of the amorphous silica shell and the underlying biomineralization process. Materials scientists and chemists are now exploiting diatoms to develop new biomimetic approaches and to create silicon-based microdevices with specific features. .COPYRGT. 2005 Elsevier Ltd. All rights reserved.

L7 ANSWER 14 OF 14 EMBASE COPYRIGHT (c) 2007 Elsevier B.V. All rights reserved on STN

ACCESSION NUMBER: 2004444353 EMBASE  
TITLE: Silica morphogenesis by alternative processing of silaffins in the diatom *Thalassiosira pseudonana*.  
AUTHOR: Poulsen N.; Kroger N.  
CORPORATE SOURCE: N. Kroger, Biochemie I, Universitat Regensburg, Universitaetsstr. 31, 93053 Regensburg, Germany. nils.kroeger@vkl.uni-regensburg.de  
SOURCE: Journal of Biological Chemistry, (8 Oct 2004) Vol. 279, No. 41, pp. 42993-42999. .  
Refs: 25  
ISSN: 0021-9258 CODEN: JBCHA3  
COUNTRY: United States  
DOCUMENT TYPE: Journal; Article  
FILE SEGMENT: 004 Microbiology  
LANGUAGE: English  
SUMMARY LANGUAGE: English  
ENTRY DATE: Entered STN: 19 Nov 2004  
Last Updated on STN: 19 Nov 2004

AB For almost 200 years scientists have been fascinated by the ornate cell walls of the diatoms. These structures are made of amorphous silica, exhibiting species-specific, mostly porous patterns in the nano- to micrometer range. Recently, from the diatom *Cylindrotheca fusiformis* unusual phosphoproteins (termed silaffins) and long chain polyamines have been identified and implicated in biosilica formation. However, analysis of the role of silaffins in morphogenesis of species-specific silica structures has so far been hampered by the difficulty of obtaining structural data from these extremely complex proteins. In the present study, the five major silaffins from the diatom *Thalassiosira pseudonana* (tpSillH, -1L, -2H, -2L, and -3) have been isolated, functionally analyzed, and structurally characterized, making use of the recently available genome data from this organism. Surprisingly, the silaffins of *T. pseudonana* and *C. fusiformis* share no sequence homology but are similar regarding amino acid composition and posttranslational modifications. Silaffins tpSillH and -2H are higher molecular mass isoforms of tpSillL and -2L, respectively, generated in vivo by alternative processing of the same precursor polypeptides. Interestingly, only tpSillH and -2H but not tpSillL and -2L induce the formation of porous silica patterns in vitro, suggesting that the alternative processing event is an important step in morphogenesis of *T. pseudonana* biosilica.

=> s silaffin and polypeptide and silica  
L5 20 SILAFFIN AND POLYPEPTIDE AND SILICA

=> l5 and immobilize  
L5 IS NOT A RECOGNIZED COMMAND  
The previous command name entered was not recognized by the system.  
For a list of commands available to you in the current file, enter  
"HELP COMMANDS" at an arrow prompt (=>).

=> s l5 and enzyme and protein  
L6 3 L5 AND ENZYME AND PROTEIN

=> l5 and protein  
L5 IS NOT A RECOGNIZED COMMAND  
The previous command name entered was not recognized by the system.  
For a list of commands available to you in the current file, enter  
"HELP COMMANDS" at an arrow prompt (=>).

=> s l5 and protein  
L7 14 L5 AND PROTEIN

=> d ibib abs l6 1-3

L6 ANSWER 1 OF 3 CAPLUS COPYRIGHT 2007 ACS on STN